

BATCH

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/626,731

DATE: 10/04/2000
 TIME: 14:32:26

Input Set : A:\626731.txt
 Output Set: N:\CRF3\10042000\I626731.raw

4 <110> APPLICANT: Co, Man Sung
 5 Vasquez, Maximiliano
 6 Carreno, Beatriz
 7 Celniker, Abbie Cheryl
 8 Collins, Mary
 9 Goldman, Samuel
 10 Gray, Gary S.
 11 Knight, Andrea
 12 O'Hara, Denise
 13 Rup, Bonita
 14 Veldman, Geertruida M.
 16 <120> TITLE OF INVENTION: HUMANIZED IMMUNOGLOBULIN REACTIVE WITH
 17 B7-2 AND METHODS OF TREATMENT THEREWITH
 20 <130> FILE REFERENCE: GI-5315
 22 <140> CURRENT APPLICATION NUMBER: 09/626,731
 23 <141> CURRENT FILING DATE: 2000-07-27
 25 <150> PRIOR APPLICATION NUMBER: 09/249,011
 26 <151> PRIOR FILING DATE: 1999-02-12
 28 <160> NUMBER OF SEQ ID NOS: 20
 30 <170> SOFTWARE: FastSEQ for Windows Version 3.0
 32 <210> SEQ ID NO: 1
 33 <211> LENGTH: 405
 34 <212> TYPE: DNA
 35 <213> ORGANISM: Murine anti-B7-2 heavy chain
 37 <220> FEATURE:
 38 <221> NAME/KEY: CDS
 39 <222> LOCATION: (1)...(405)
 40 <223> OTHER INFORMATION:
 47 <400> SEQUENCE: 1 48
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 49 Met Gly Trp Asn Cys Ile Ile Phe Phe Leu Val Thr Thr Ala Thr Gly 15 96
 50 1 5 10 15
 52 gtg cac tcc cag gtc cag ctg cag tct ggg cct gag ctg gtg agg 30
 53 Val His Ser Gln Val Gln Leu Gln Ser Gly Pro Glu Leu Val Arg 30
 54 20 25 30
 56 cct ggg gaa tca gtg aag att tcc tgc aag ggt tcc ggc tac aca ttc 144
 57 Pro Gly Glu Ser Val Lys Ile Ser Cys Lys Gly Ser Gly Tyr Thr Phe 45
 58 35 40 45
 60 act gat tat gct ata cag tgg gtg aag cag agt cat gca aag agt cta 192
 61 Thr Asp Tyr Ala Ile Gln Trp Val Lys Gln Ser His Ala Lys Ser Leu 55 60
 62 50 55 60
 64 gag tgg att gga gtt att aat att tac tat gat aat aca aac tac aac 240
 65 Glu Trp Ile Gly Val Ile Asn Ile Tyr Tyr Asp Asn Thr Asn Tyr Asn 75 80
 66 65 70 75 80
 68 cag aag ttt aag ggc aag gcc aca atg act gta gac aaa tcc tcc agc 288
 69 Gln Lys Phe Lys Gly Lys Ala Thr Met Thr Val Asp Lys Ser Ser Ser 85 90 95
 70 75 80 85 90 95

ENTERED

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72	aca gcc tat atg gaa ctt gcc aga ttg aca tct gag gat tct gcc atc	336
73	Thr Ala Tyr Met Glu Leu Ala Arg Leu Thr Ser Glu Asp Ser Ala Ile	
74	100 105 110	
76	tat tac tgt gca aga gcg gcc tgg tat atg gac tac tgg ggt caa gga	384
77	Tyr Tyr Cys Ala Arg Ala Ala Trp Tyr Met Asp Tyr Trp Gly Gln Gly	
78	115 120 125	
80	acc tca gtc acc gtc tcc tca	405
81	Thr Ser Val Thr Val Ser Ser	
82	130 135	
85	<210> SEQ ID NO: 2	
86	<211> LENGTH: 135	
87	<212> TYPE: PRT	
88	<213> ORGANISM: Murine anti-B7-2 heavy chain	
90	<400> SEQUENCE: 2	
91	Met Gly Trp Asn Cys Ile Ile Phe Phe Leu Val Thr Thr Ala Thr Gly	
92	1 5 10 15	
93	Val His Ser Gln Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Arg	
94	20 25 30	
95	Pro Gly Glu Ser Val Lys Ile Ser Cys Lys Gly Ser Gly Tyr Thr Phe	
96	35 40 45	
97	Thr Asp Tyr Ala Ile Gln Trp Val Lys Gln Ser His Ala Lys Ser Leu	
98	50 55 60	
99	Glu Trp Ile Gly Val Ile Asn Ile Tyr Tyr Asp Asn Thr Asn Tyr Asn	
100	65 70 75 80	
101	Gln Lys Phe Lys Gly Lys Ala Thr Met Thr Val Asp Lys Ser Ser Ser	
102	85 90 95	
103	Thr Ala Tyr Met Glu Leu Ala Arg Ile Thr Ser Glu Asp Ser Ala Ile	
104	100 105 110	
105	Tyr Tyr Cys Ala Arg Ala Ala Trp Tyr Met Asp Tyr Trp Gly Gln Gly	
106	115 120 125	
107	Thr Ser Val Thr Val Ser Ser	
108	130 135	
110	<210> SEQ ID NO: 3	
111	<211> LENGTH: 396	
112	<212> TYPE: DNA	
113	<213> ORGANISM: Murine anti-B7-2 light chain	
115	<220> FEATURE:	
116	<221> NAME/KEY: CDS	
117	<222> LOCATION: (1)...(396)	
119	<400> SEQUENCE: 3	
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121	Met Asp Ser Gln Ala Gln Val Leu Ile Leu Leu Leu Trp Val Ser	
122	1 5 10 15	
124	ggc acc tgt ggg gac att gtg ctg tca cag tct cca tcc tcc ctg gct	96
125	Gly Thr Cys Gly Asp Ile Val Leu Ser Gln Ser Pro Ser Ser Leu Ala	
126	20 25 30	
128	gtg tca gca gga gag aag gtc act atg agc tgc aaa tcc agt cag agt	144
129	Val Ser Ala Gly Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser	
130	35 40 45	

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132	ctg ctc aac agt aga acc cga gag aac tac ttg gct tgg tac cag cag	192
133	Leu Leu Asn Ser Arg Thr Arg Glu Asn Tyr Leu Ala Trp Tyr Gln Gln	
134	50 55 60	
136	aaa cca ggg cag tct cct aaa ctg ctg atc tac tgg gca tcc act agg	240
137	Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg	
138	65 70 75 80	
140	gaa tct ggg gtc cct gat cgc ttc aca ggc agt gga tct ggg aca gat	288
141	Glu Ser Gly Val Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp	
142	85 90 95	
144	ttc act ctc acc atc agc agt gtg cag gct gaa gac ctg gca gtt tat	336
145	Phe Thr Leu Thr Ile Ser Ser Val Gln Ala Glu Asp Leu Ala Val Tyr	
146	100 105 110	
148	tac tgc acg caa tct tat aat ctt tac acg ttc gga ggg ggg acc aag	384
149	Tyr Cys Thr Gln Ser Tyr Asn Leu Tyr Thr Phe Gly Gly Thr Lys	
150	115 120 125	
152	ctg gaa ata aaa	396
153	Leu Glu Ile Lys	
154	130	
157	<210> SEQ ID NO: 4	
158	<211> LENGTH: 132	
159	<212> TYPE: PRT	
160	<213> ORGANISM: Murine anti-B7-2 light chain	
162	<400> SEQUENCE: 4	
163	Met Asp Ser Gln Ala Gln Val Leu Ile Leu Leu Leu Leu Trp Val Ser	
164	1 5 10 15	
165	Gly Thr Cys Gly Asp Ile Val Leu Ser Gln Ser Pro Ser Ser Leu Ala	
166	20 25 30	
167	Val Ser Ala Gly Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser	
168	35 40 45	
169	Leu Leu Asn Ser Arg Thr Arg Glu Asn Tyr Leu Ala Trp Tyr Gln Gln	
170	50 55 60	
171	Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg	
172	65 70 75 80	
173	Glu Ser Gly Val Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp	
174	85 90 95	
175	Phe Thr Leu Thr Ile Ser Ser Val Gln Ala Glu Asp Leu Ala Val Tyr	
176	100 105 110	
177	Tyr Cys Thr Gln Ser Tyr Asn Leu Tyr Thr Phe Gly Gly Thr Lys	
178	115 120 125	
179	Leu Glu Ile Lys	
180	130	
182	<210> SEQ ID NO: 5	
183	<211> LENGTH: 405	
184	<212> TYPE: DNA	
185	<213> ORGANISM: Humanized murine anti-human B7-2 heavy chain	
187	<220> FEATURE:	
188	<221> NAME/KEY: CDS	
189	<222> LOCATION: (1)...(405)	
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192	atg ggt tgg aac tgt atc atc ttc ttt ctg gtt acc aca gct aca ggt	48
193	Met Gly Trp Asn Cys Ile Ile Phe Phe Leu Val Thr Thr Ala Thr Gly	
194	1 5 10 15	
196	gtg cac tcc cag gtc cag ctg gtg cag tct ggg gct gag gtg aag aag	96
197	Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys	
198	20 25 30	
200	cct ggg agc tca gtg aag gtg tcc tgc aaa gct tcc ggc tac aca ttc	144
201	Pro Gly Ser Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe	
202	35 40 45	
204	act gat tat gct ata cag tgg gtg aga cag gct cct gga cag ggc ctc	192
205	Thr Asp Tyr Ala Ile Gln Trp Val Arg Gln Ala Pro Gly Gln Gly Leu	
206	50 55 60	
208	gag tgg att gga gtt att aat att tac tat gat aat aca aac tac aac	240
209	Glu Trp Ile Gly Val Ile Asn Ile Tyr Tyr Asn Thr Asn Tyr Asn	
210	65 70 75 80	
212	cag aag ttt aag ggc aag gcc aca atg act gta gac aag tcg acg acg	288
213	Gln Lys Phe Lys Gly Lys Ala Thr Met Thr Val Asp Lys Ser Thr Ser	
214	85 90 95	
216	aca gcc tat atg gaa ctt agt tct ttg aga tct gag gat acg gcc gtt	336
217	Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val	
218	100 105 110	
220	tat tac tgt gca aga gcg gcc tgg tat atg gac tac tgg ggt caa ggt	384
221	Tyr Tyr Cys Ala Arg Ala Ala Trp Tyr Met Asp Tyr Trp Gly Gln Gly	
222	115 120 125	
224	acc ctt gtc acc gtc tcc tca	405
225	Thr Leu Val Thr Val Ser Ser	
226	130 135	
229	<210> SEQ ID NO: 6	
230	<211> LENGTH: 135	
231	<212> TYPE: PRT	
232	<213> ORGANISM: Humanized murine anti-human B7-2 heavy chain	
234	<400> SEQUENCE: 6	
235	Met Gly Trp Asn Cys Ile Ile Phe Phe Leu Val Thr Thr Ala Thr Gly	
236	1 5 10 15	
237	Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys	
238	20 25 30	
239	Pro Gly Ser Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe	
240	35 40 45	
241	Thr Asp Tyr Ala Ile Gln Trp Val Arg Gln Ala Pro Gly Gln Gly Leu	
242	50 55 60	
243	Glu Trp Ile Gly Val Ile Asn Ile Tyr Tyr Asp Asn Thr Asn Tyr Asn	
244	65 70 75 80	
245	Gln Lys Phe Lys Gly Lys Ala Thr Met Thr Val Asp Lys Ser Thr Ser	
246	85 90 95	
247	Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val	
248	100 105 110	
249	Tyr Tyr Cys Ala Arg Ala Ala Trp Tyr Met Asp Tyr Trp Gly Gln Gly	
250	115 120 125	
251	Thr Leu Val Thr Val Ser Ser	

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255 <211>	LENGTH: 396		
256 <212>	TYPE: DNA		
257 <213>	ORGANISM: Humanized murine anti-human B7-2 light chain		
259 <220>	FEATURE:		
260 <221>	NAME/KEY: CDS		
261 <222>	LOCATION: (1)...(396)		
263 <400>	SEQUENCE: 7		
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266 1 5 10 15			
268 ggc acc tgt ggg gac att gtg ctg aca cag tct cca gat tcc ctg gct		96	
269 Gly Thr Cys Gly Asp Ile Val Leu Thr Gln Ser Pro Asp Ser Leu Ala			
270 20 25 30			
272 gta agc tta gga gag agg gcc act att agc tgc aaa tcc agt cag agt		144	
273 Val Ser Leu Gly Glu Arg Ala Thr Ile Ser Cys Lys Ser Ser Gln Ser			
274 35 40 45			
276 ctg ctc aac agt aga acc cga gag aac tac ttg gct tgg tac cag cag		192	
277 Leu Leu Asn Ser Arg Thr Arg Glu Asn Tyr Leu Ala Trp Tyr Gln Gln			
278 50 55 60			
280 aaa cca ggg cag cct cct aaa ctg ctg atc tac ttg gca tcc act agg		240	
281 Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg			
282 65 70 75 80			
284 gaa tct ggg gtc cct gat cgc ttc agt ggc agt gga tct ggg aca gat		288	
285 Glu Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp			
286 85 90 95			
288 ttc act ctc acc atc agc agt ctg cag gct gaa gac gtc gca gtt tat		336	
289 Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr			
290 100 105 110			
292 tac tgc acg caa tct tat aat ctt tac acg ttc gga cag ggg acc aag		384	
293 Tyr Cys Thr Gln Ser Tyr Asn Leu Tyr Thr Phe Gly Gln Gly Thr Lys			
294 115 120 125			
296 gtg gaa ata aaa		396	
297 Val Glu Ile Lys			
298 130			
301 <210>	SEQ ID NO: 8		
302 <211>	LENGTH: 132		
303 <212>	TYPE: PRT		
304 <213>	ORGANISM: Humanized murine anti-human B7-2 light chain		
306 <400>	SEQUENCE: 8		
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309 Gly Thr Cys Gly Asp Ile Val Leu Thr Gln Ser Pro Asp Ser Leu Ala			
310 20 25 30			
311 Val Ser Leu Gly Glu Arg Ala Thr Ile Ser Cys Lys Ser Ser Gln Ser			
312 35 40 45			
313 Leu Leu Asn Ser Arg Thr Arg Glu Asn Tyr Leu Ala Trp Tyr Gln Gln			
314 50 55 60			

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